

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/540,494

Source: PCT/10

Date Processed by STIC: 7/11/05

ENTERED



PCT

RAW SEQUENCE LISTING

DATE: 07/11/2005

PATENT APPLICATION: US/10/540,494

TIME: 15:08:42

Input Set : A:\63628 Sequence Listing.txt

Output Set: N:\CRF4\07112005\J540494.raw

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3 <110> APPLICANT: Takeda Pharmaceutical Company Limited
5 <120> TITLE OF INVENTION: Metastatin Derivatives And Its Use
7 <130> FILE REFERENCE: G05-0018
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/540,494
C--> 9 <141> CURRENT FILING DATE: 2005-06-23
9 <150> PRIOR APPLICATION NUMBER: PCT/JP2003/016978
10 <151> PRIOR FILING DATE: 2003-12-26
12 <150> PRIOR APPLICATION NUMBER: JP 2002-377179
13 <151> PRIOR FILING DATE: 2002-12-26
15 <160> NUMBER OF SEQ ID NOS: 22
17 <210> SEQ ID NO: 1
18 <211> LENGTH: 54
19 <212> TYPE: PRT
20 <213> ORGANISM: Homo sapiens
22 <400> SEQUENCE: 1
23 Gly Thr Ser Leu Ser Pro Pro Pro Glu Ser Ser Gly Ser Arg Gln Gln
24 1 5 10 15
25 Pro Gly Leu Ser Ala Pro His Ser Arg Gln Ile Pro Ala Pro Gln Gly
26 20 25 30
27 Ala Val Leu Val Gln Arg Glu Lys Asp Leu Pro Asn Tyr Asn Trp Asn
28 35 40 45
29 Ser Phe Gly Leu Arg Phe
30 50
32 <210> SEQ ID NO: 2
33 <211> LENGTH: 162
34 <212> TYPE: DNA
35 <213> ORGANISM: Homo sapiens
37 <400> SEQUENCE: 2
38 ggtacttctc tgtctccgcc gccggaatct tctggttctc gtcagcagcc ggtctgtct 60
39 gctccgcact ctgctcagat cccggctccg caggggtgctg ttctggttca gcgtgaaaaa 120
40 gacctgccga actacaactg gaactctttc ggtctgcgtt tc 162
42 <210> SEQ ID NO: 3
43 <211> LENGTH: 152
44 <212> TYPE: PRT
45 <213> ORGANISM: Mus musculus
47 <400> SEQUENCE: 3
48 Met Tyr Leu Arg Phe Gly Val Asp Val Cys Ser Leu Ser Pro Trp Lys
49 5 10 15
50 Glu Thr Val Asp Leu Pro Leu Pro Pro Arg Met Ile Ser Met Ala Ser
51 20 25 30
52 Trp Gln Leu Leu Leu Leu Cys Val Ala Thr Tyr Gly Glu Pro Leu
53 35 40 45
54 Ala Lys Val Ala Pro Gly Ser Thr Gly Gln Gln Ser Gly Pro Gln Glu

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55      50      55      60
56 Leu Val Asn Ala Trp Glu Lys Glu Ser Arg Tyr Ala Glu Ser Lys Pro
57 65      70      75      80
58 Gly Ser Ala Gly Leu Arg Ala Arg Arg Ser Ser Pro Cys Pro Pro Val
59      85      90      95
60 Glu Gly Pro Ala Gly Arg Gln Arg Pro Leu Cys Ala Ser Arg Ser Arg
61      100      105      110
62 Leu Ile Pro Ala Pro Arg Gly Ala Val Leu Val Gln Arg Glu Lys Asp
63      115      120      125
64 Leu Ser Thr Tyr Asn Trp Asn Ser Phe Gly Leu Arg Tyr Gly Arg Arg
65      130      135      140
66 Gln Ala Ala Arg Ala Ala Arg Gly
67 145      150
69 <210> SEQ ID NO: 4
70 <211> LENGTH: 456
71 <212> TYPE: DNA
72 <213> ORGANISM: Mus musculus
74 <400> SEQUENCE: 4
75 atgtatctga gatttggcgt tgatgtctgc agcctgagtc cctggaagga gactgtagac 60
76 ctgcccccttc ctcccagaat gatctcaatg gcttcttggc agctgctgct tctcctctgt 120
77 gtcgccacct atggggagcc gctggcaaaa gtgaagcctg gatccacagg ccagcagtcc 180
78 ggaccccagg aactcgttaa tgcctgggaa aaggaatcgc ggtatgcaga gagcaagcct 240
79 gggctctgcag ggctgcgcgc tcgtaggtcg tcgccatgcc cgccggttga gggccccgcg 300
80 gggcgccagc ggccccctgtg tgcctcccgc agtcgcctga tccctgcgcc ccgcggagcg 360
81 gtgctggtgc agcgggagaa ggacctgtcc acctacaact ggaactcctt cggcctgcgc 420
82 tacggcagga ggcaggcggc gcgggcagca cggggc 456
84 <210> SEQ ID NO: 5
85 <211> LENGTH: 156
86 <212> TYPE: PRT
87 <213> ORGANISM: Mus musculus
89 <400> SEQUENCE: 5
90 Met Tyr Leu Arg Phe Gly Val Asp Val Cys Ser Leu Ser Pro Trp Lys
91      5      10      15
92 Glu Thr Val Asp Leu Pro Leu Pro Pro Arg Met Ile Ser Met Ala Ser
93      20      25      30
94 Trp Gln Leu Leu Leu Leu Leu Cys Val Ala Thr Tyr Gly Glu Pro Leu
95      35      40      45
96 Ala Lys Val Ala Pro Leu Val Lys Pro Gly Ser Thr Gly Gln Gln Ser
97      50      55      60
98 Gly Pro Gln Glu Leu Val Asn Ala Trp Glu Lys Glu Ser Arg Tyr Ala
99 65      70      75      80
100 Glu Ser Lys Pro Gly Ser Ala Gly Leu Arg Ala Arg Arg Ser Ser Pro
101      85      90      95
102 Cys Pro Pro Val Glu Gly Pro Ala Gly Arg Gln Arg Pro Leu Cys Ala
103      100      105      110
104 Ser Arg Ser Arg Leu Ile Pro Ala Pro Arg Gly Ala Val Leu Val Gln
105      115      120      125
106 Arg Glu Lys Asp Leu Ser Thr Tyr Asn Trp Asn Ser Phe Gly Leu Arg
107      130      135      140

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108 Tyr Gly Arg Arg Gln Ala Ala Arg Ala Ala Arg Gly
109 145          150          155
111 <210> SEQ ID NO: 6
112 <211> LENGTH: 468
113 <212> TYPE: DNA
114 <213> ORGANISM: Mus musculus
116 <400> SEQUENCE: 6
117 atgtatctga gatttggcgt tgatgtctgc agcctgagtc cctggaagga gactgtagac * 60
118 ctgccccttc ctcccagaat gatctcaatg gcttcttggc agctgctgct tctcctctgt 120
119 gtcgccacct atggggagcc gctggcaaaa gtggcacctt tggatgaagcc tggatccaca 180
120 ggccagcagt ccggacccca ggaactcggt aatgcctggg aaaaggaatc gcggtatgca 240
121 gagagcaagc ctgggtctgc agggctgcgc gctcgtaggt cgtcgccatg cccgccggtt 300
122 gagggccccc cggggcgcca gcggccctg tgtgcctccc gcagtcgcct gatccctgcg 360
123 ccccgcgag cggtgctggt gcagcgggag aaggacctgt ccacctaca ctggaactcc 420
124 ttcggcctgc gctacggcag gaggcaggcg gcgcgggcag cacggggc 468
126 <210> SEQ ID NO: 7
127 <211> LENGTH: 130
128 <212> TYPE: PRT
129 <213> ORGANISM: Rattus sp.
131 <400> SEQUENCE: 7
132 Met Thr Ser Leu Ala Ser Trp Gln Leu Leu Leu Leu Leu Cys Val Ala
133          5          10          15
134 Ser Phe Gly Glu Pro Leu Ala Lys Met Ala Pro Val Val Asn Pro Glu
135          20          25          30
136 Pro Thr Gly Gln Gln Ser Gly Pro Gln Glu Leu Val Asn Ala Trp Gln
137          35          40          45
138 Lys Gly Pro Arg Tyr Ala Glu Ser Lys Pro Gly Ala Ala Gly Leu Arg
139          50          55          60
140 Ala Arg Arg Thr Ser Pro Cys Pro Pro Val Glu Asn Pro Thr Gly His
141          65          70          75          80
142 Gln Arg Pro Pro Cys Ala Thr Arg Ser Arg Leu Ile Pro Ala Pro Arg
143          85          90          95
144 Gly Ser Val Leu Val Gln Arg Glu Lys Asp Met Ser Ala Tyr Asn Trp
145          100          105          110
146 Asn Ser Phe Gly Leu Arg Tyr Gly Arg Arg Gln Val Ala Arg Ala Ala
147          115          120          125
148 Arg Gly
149          130
151 <210> SEQ ID NO: 8
152 <211> LENGTH: 390
153 <212> TYPE: DNA
154 <213> ORGANISM: Rattus sp.
156 <400> SEQUENCE: 8
157 atgacctgc tggttcttg gcagctgctg cttctcctct gtgtggcctc ttttggggag 60
158 ccactggcaa aaatggcacc tgttgtgaac cctgaaccca caggccaaca gtccggaccc 120
159 caggaactcg ttaatgcctg gcaaaagggc ccgcggtatg cagagagcaa gcctggggct 180
160 gcaggactgc gcgctgcgcg aacatcgcca tgcccgcggg tggagaaccc cacggggcac 240
161 cagcggcccc cgtgtgccac ccgcagtcgc ctgatccctg cgccccgcgg atcggtgctg 300
162 gtgcagcgcg agaaggacat gtcagcctac aactggaact cctttggcct gcgctacggc 360

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163 aggaggcagg tggcgcgggc ggcacggggc 390
165 <210> SEQ ID NO: 9
166 <211> LENGTH: 398
167 <212> TYPE: PRT
168 <213> ORGANISM: Homo sapiens
170 <400> SEQUENCE: 9
171 Met His Thr Val Ala Thr Ser Gly Pro Asn Ala Ser Trp Gly Ala Pro
172           5           10           15
173 Ala Asn Ala Ser Gly Cys Pro Gly Cys Gly Ala Asn Ala Ser Asp Gly
174           20           25           30
175 Pro Val Pro Ser Pro Arg Ala Val Asp Ala Trp Leu Val Pro Leu Phe
176           35           40           45
177 Phe Ala Ala Leu Met Leu Leu Gly Leu Val Gly Asn Ser Leu Val Ile
178           50           55           60
179 Tyr Val Ile Cys Arg His Lys Pro Met Arg Thr Val Thr Asn Phe Tyr
180           65           70           75           80
181 Ile Ala Asn Leu Ala Ala Thr Asp Val Thr Phe Leu Leu Cys Cys Val
182           85           90           95
183 Pro Phe Thr Ala Leu Leu Tyr Pro Leu Pro Gly Trp Val Leu Gly Asp
184           100          105          110
185 Phe Met Cys Lys Phe Val Asn Tyr Ile Gln Gln Val Ser Val Gln Ala
186           115          120          125
187 Thr Cys Ala Thr Leu Thr Ala Met Ser Val Asp Arg Trp Tyr Val Thr
188           130          135          140
189 Val Phe Pro Leu Arg Ala Leu His Arg Arg Thr Pro Arg Leu Ala Leu
190           145          150          155          160
191 Ala Val Ser Leu Ser Ile Trp Val Gly Ser Ala Ala Val Ser Ala Pro
192           165          170          175
193 Val Leu Ala Leu His Arg Leu Ser Pro Gly Pro Arg Ala Tyr Cys Ser
194           180          185          190
195 Glu Ala Phe Pro Ser Arg Ala Leu Glu Arg Ala Phe Ala Leu Tyr Asn
196           195          200          205
197 Leu Leu Ala Leu Tyr Leu Leu Pro Leu Leu Ala Thr Cys Ala Cys Tyr
198           210          215          220
199 Ala Ala Met Leu Arg His Leu Gly Arg Val Ala Val Arg Pro Ala Pro
200           225          230          235          240
201 Ala Asp Ser Ala Leu Gln Gly Gln Val Leu Ala Glu Arg Ala Gly Ala
202           245          250          255
203 Val Arg Ala Lys Val Ser Arg Leu Val Ala Ala Val Val Leu Leu Phe
204           260          265          270
205 Ala Ala Cys Trp Gly Pro Ile Gln Leu Phe Leu Val Leu Gln Ala Leu
206           275          280          285
207 Gly Pro Ala Gly Ser Trp His Pro Arg Ser Tyr Ala Ala Tyr Ala Leu
208           290          295          300
209 Lys Thr Trp Ala His Cys Met Ser Tyr Ser Asn Ser Ala Leu Asn Pro
210           305          310          315          320
211 Leu Leu Tyr Ala Phe Leu Gly Ser His Phe Arg Gln Ala Phe Arg Arg
212           325          330          335
213 Val Cys Pro Cys Ala Pro Arg Arg Pro Arg Arg Pro Arg Arg Pro Gly

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214          340          345          350
215 Pro Ser Asp Pro Ala Ala Pro His Ala Glu Leu His Arg Leu Gly Ser
216          355          360          365
217 His Pro Ala Pro Ala Arg Ala Gln Lys Pro Gly Ser Ser Gly Leu Ala
218          370          375          380
219 Ala Arg Gly Leu Cys Val Leu Gly Glu Asp Asn Ala Pro Leu
220 385          390          395
222 <210> SEQ ID NO: 10
223 <211> LENGTH: 1194
224 <212> TYPE: DNA
225 <213> ORGANISM: Homo sapiens
227 <400> SEQUENCE: 10
228 atgcacaccg tggctacgtc cggacccaac gcgtcctggg gggcaccggc caacgcctcc 60
229 ggctgcccgg gctgtggcgc caacgcctcg gacggcccag tcccttcgcc gcggggccgtg 120
230 gacgcctggc tcgtgccgct cttcttcgcg gcgtgatgc tgcgtggcct ggtggggaac 180
231 tcgctgggtca tctacgtcat ctgccgccac aagccgatgc ggaccgtgac caacttctac 240
232 atcgccaacc tggcgcccac ggacgtgacc ttctcctgt gctgcgtccc cttcacggcc 300
233 ctgctgtacc cgctgcccgg ctgggtgctg ggcgacttca tgtgcaagtt cgtcaactac 360
234 atccagcagg tctcggtgca ggccacgtgt gccactctga ccgccatgag tgtggaccgc 420
235 tggtagctga cgggtgtccc gttgcgcgcc ctgcaccgcc gcacgccccg cctggcgctg 480
236 gctgtcagcc tcagcatctg ggtaggctct gcggcggtgt ctgcgcgggt gctcgccctg 540
237 caccgcctgt caccggggcc gcgcgcctac tgcagtgagg ccttccccag ccgcgcctg 600
238 gagcgcgctt tcgactgta caacctgctg gcgctgtacc tgctgccgct gctcgccacc 660
239 tgcgcctgct atgcggccat gctgcgccac ctgggcccgg tcgccgtgcg ccccgcgccc 720
240 gccgatagcg ccttcgaggg gcaggtgctg gcagagcgcg caggcgccgt gcggggccaa 780
241 gtctcgcggc tgggtggcgg cgtggctcct ctcttcgccg cctgctgggg ccccatccag 840
242 ctgttcctgg tctgcaggg cctggggccc gcgggctcct ggcaaccacg cagctacgcc 900
243 gcctacgcgc ttaagacctg ggctcactgc atgtcctaca gcaactccgc gctgaacccg 960
244 ctgctctacg ctttcctggg ctgcgacttc cgacaggcct tccgcccgtg ctgcccctgc 1020
245 gcgcgcgcgc gccccgcgcg cccccgcgcg cccggaccct cggaccccgc agccccacac 1080
246 gcggagctgc accgcctggg gtcccaccgc gccccgcga gggcgagaa gccagggagc 1140
247 agtgggctgg ccgcgcgcgg gctgtgcgtc ctgggggagg acaacgcccc tctc 1194
249 <210> SEQ ID NO: 11
250 <211> LENGTH: 396
251 <212> TYPE: PRT
252 <213> ORGANISM: Rattus sp.
254 <400> SEQUENCE: 11
255 Met Ala Ala Glu Ala Thr Leu Gly Pro Asn Val Ser Trp Trp Ala Pro
256          5          10          15
257 Ser Asn Ala Ser Gly Cys Pro Gly Cys Gly Val Asn Ala Ser Asp Gly
258          20          25          30
259 Pro Gly Ser Ala Pro Arg Pro Leu Asp Ala Trp Leu Val Pro Leu Phe
260          35          40          45
261 Phe Ala Ala Leu Met Leu Leu Gly Leu Val Gly Asn Ser Leu Val Ile
262          50          55          60
263 Phe Val Ile Cys Arg His Lys His Met Gln Thr Val Thr Asn Phe Tyr
264 65          70          75          80
265 Ile Ala Asn Leu Ala Ala Thr Asp Val Thr Phe Leu Leu Cys Cys Val
266          85          90          95

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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/540,494

DATE: 07/11/2005
TIME: 15:08:43

Input Set : A:\63628 Sequence Listing.txt
Output Set: N:\CRF4\07112005\J540494.raw

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:15,16,17,18

VERIFICATION SUMMARY

DATE: 07/11/2005

PATENT APPLICATION: US/10/540,494

TIME: 15:08:43

Input Set : A:\63628 Sequence Listing.txt

Output Set: N:\CRF4\07112005\J540494.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application No

L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date